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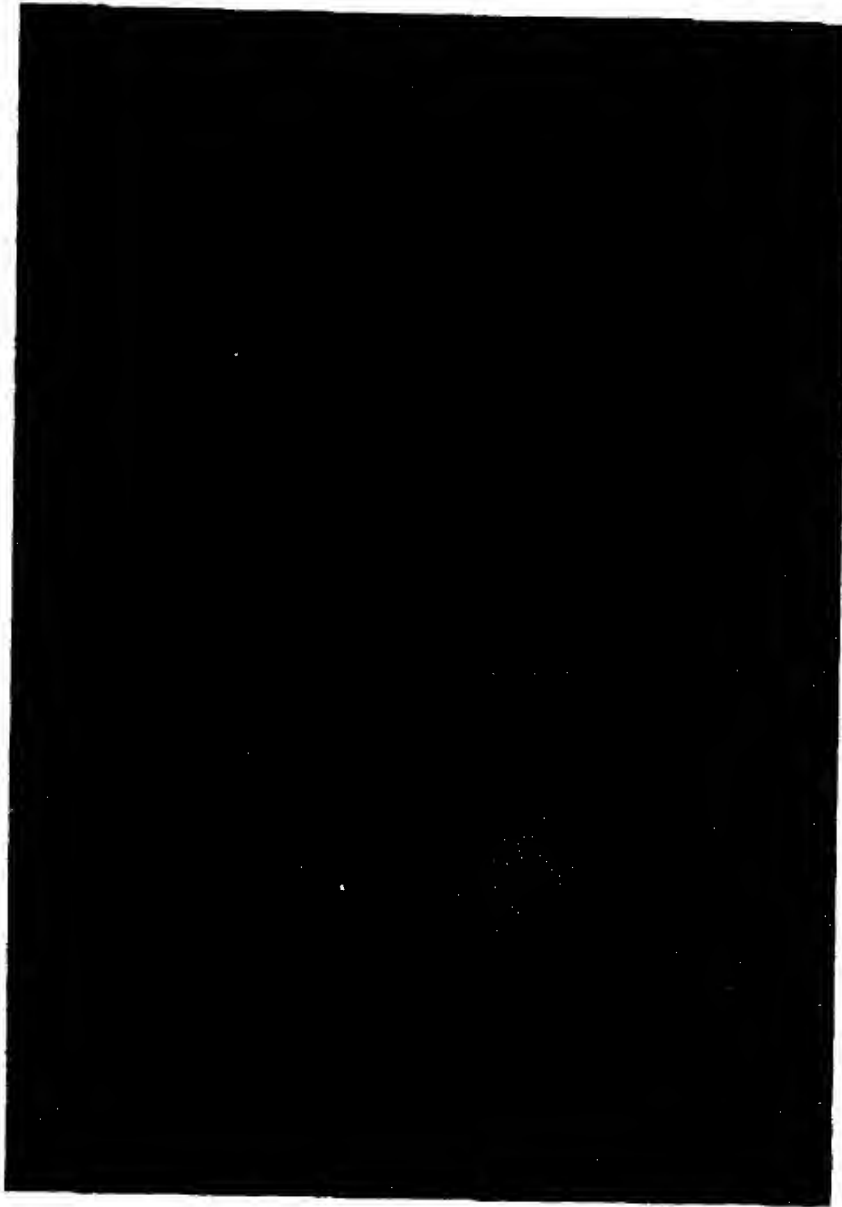


Fig. 1

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[illegible]

**Fig. 2a**



|                |     |                                 |                |   |                      |                 |               |
|----------------|-----|---------------------------------|----------------|---|----------------------|-----------------|---------------|
| HS-HMC-Dis3    | 183 | AIQEGIPAPTCEEYVKTLT             | ----           | ANPELLIDRLACLSEEGNEIES                        | ----                 | GKILFSEHPLPSKL  |               |
| MM-hmc-Dis3    | 183 | AVQEGIPAPTCEEYVKSLT             | ----           | ANPELLIDRLAYLSDEMNEIES                        | ----                 | GKILFSEHPLPSKL  |               |
| CE-Q17632-Dis3 | 191 | HYQH----                        | VMYLKEYIQNLEDP | ----  | GKQALLDQMAAYESSGNENE | ----            | KQIFDEVLSHDRI |
| SP-P37202-Dis3 | 199 | AAEQGIQVSTLKDYVQYLP             | ----           | DSEILLDMVSAIADAIASKEQVES                      | ----                 | GTNNVVELHWSMSRL |               |
| SC-Q08162-Dis3 | 207 | EVESNIITKSLVQYIELLP             | ----           | NADDIRDSIP-QMDSFDKDLERDT                      | ----                 | FSDFTTPEXYSTARV |               |
| MM-LOBO        | 107 | GDLVVVKLLPEDQKAVKP              | ----           | E----   | SNDKEIETYEADIPE      | ----            | EGCGHHPL      |
| HS-LOBO        | 0   |                                 |                |   |                      |                 |               |
| CE-Q09568-LOBO | 92  | GDVVAVKIKPKEDWLVNYV             | ----           | E----   | YVKWMAEH             | ----            |               |
| SP-Z99259      | 143 | NSAK--REKNNSHQVEADT             | ----           | NN--ATENVSSNAKKS                              | ----                 | VYPLYYDSATV     |               |
| SC-P24276-SSD1 | 280 | PQQQLSPFRHRGSNSRDYNSFNTLEPPAI   | ----           | FQQCHKHRA SNSSVHSFSSQGNNGGGRKSLFAPYLPQANI     | ----                 |                 |               |
| SC-P39112      | 206 | NRVLLRIPHKLPAGIHSLI             | ----           | QPESHKHLPIGTVKNFNSQTN                         | ----                 | ILPIVARQLITSRY  |               |
|                | 281 | 291                             | 301            | 311   | 321                  | 331             | 341           |
| HS-HMC-Dis3    | 237 | QQGIKSGTYLQGTFRASRENYLEATV      | ----           | WIHGDNENKEIILQGLKHLNRAVHEDIVAVELLPKS-QMVAAPS  |                      |                 |               |
| MM-hmc-Dis3    | 237 | QQGIKSGSYLQGTFRASRENYLEATV      | ----           | WIHGDKCEEKEIILQGIKHLNRAVHEDIVAVELLPRS-QMVAAPS |                      |                 |               |
| CE-Q17632-Dis3 | 242 | MEGIASGTIKRGNFVSRENYREATVIID    | ----           | DQTSWFITG-NNCNRAVNGDTVAVQLLPED-QMTAPE         |                      |                 |               |
| SP-P37202-Dis3 | 257 | LACIKNGEVHKGLINISTYNYLEGSVVVP   | ----           | GYNKPVLVSGRENLRNAVQGDIVCICILPQD-QMKTAE        |                      |                 |               |
| SC-Q08162-Dis3 | 264 | MGGLKNGVLVYQGNIIQISEYNPLEGSVSLP | ----           | RFSEKPVLLVGGKNLNRAFNGDQVIVELLPQS-EWKAPS       |                      |                 |               |

Fig. 2c

**Fig. 2d**

**Fig. 2e**



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|                |     |  |
|----------------|-----|--|
| HS-HMC-Dis3    | 522 | -----SARRGTTTVYLCEKRIDMVPELLSS--NLCSLKCDVDRLAFSCIWEMNHNA-----EILK  |
| MM-hmc-Dis3    | 522 | -----SARRGTTTVYLCEKRIDMVPELLSS--NLCSLRSNVDRDLAFSCIWEMNHNA-----EILK |
| CE-Q17632-Dis3 | 534 | -----AALRGTTTVYLCDDRRIDMLPCLLSS--NLCSLRGEEREYAFSCIWTMTSSA-----DIQS |
| SP-P37202-Dis3 | 551 | -----AASRGTTTVYLVDKRIDMLPMLLGT--DLCSLRPYVERFAFSCIWEMDENA-----NIK   |
| SC-Q08162-Dis3 | 586 | -----GAARGTSVYLVDKRIDMLPMLLGT--DLCSLKPYVDRFAFSCIWELDDSA-----NIVN   |
| MM-LOBO        | 424 | -----AAERATSVYLVQKVVPMLPRLLCE--ELCSLNPMTDKLTFSVIWKLTPEG-----KILE   |
| HS-LOBO        | 51  | -----AAERATSVYLVQKVVPMLPRLLCE--ELCSLNPMSDXLTFSVIWTLTPEG-----KILD   |

Fig. 2g



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|                |     |   |     |     |     |     |     |     |     |
|----------------|-----|---|-----|-----|-----|-----|-----|-----|-----|
| CE-Q09568-LOBO | 363 | -----ASRGNSTYLSQTVIPMLPRILCE--QLCSINPGVDRLSFSVTFKMSYEA-----ELYD         | 771 | 781 | 791 | 801 | 811 | 821 | 831 |
| SP-Z99259      | 496 | -----AASRATTVLVQKAIPMLPPLICE--RLCSINPNVERLAFSVFWKLDNGK-----EIGK         |     |     |     |     |     |     |     |
| SC-P24276-SSD1 | 746 | -----ARKRSSAVFMPQKLVNLLPQSFN---DELSLAPGKESATLSVVYTLDSSTL-----RIKS       |     |     |     |     |     |     |     |
| SC-P39112      | 578 | GISTDIILNVALKRSFTTFLPDTVPVMPLPQSICHLSDLGKQGORTKTSFSVDVKITSKCSGKSIEIMYDS |     |     |     |     |     |     |     |
|                |     | ----- ----- ----- ----- ----- ----- ----- ----- -----                   |     |     |     |     |     |     |     |
|                |     | 771   | 781 | 791 | 801 | 811 | 821 | 831 |     |
| HS-HMC-Dis3    | 574 | TKFTKSVINSKASL-YAEAQLRIDSANMND-----DITTSRLGCLNKLAKILKKRRIENGA           |     |     |     |     |     |     |     |
| MM-hmc-Dis3    | 574 | TRFTKSVINSKASL-YAEAQMRIDSAAMND-----DITTSRLGCLNQLAKILKKGRIEKGA           |     |     |     |     |     |     |     |
| CE-Q17632-Dis3 | 586 | VKYHKSLIKSKAALTYEKAQEIIDDPKEQN-----DVALGLRGLMKLSKVLNARRTGNGA            |     |     |     |     |     |     |     |
| SP-P37202-Dis3 | 603 | VHFTKSVIASKEAFSYADAQARIDDKQMDD-----PLTQGMRVLLKLSKILKQKRMDEGA            |     |     |     |     |     |     |     |
| SC-Q08162-Dis3 | 638 | VNFMKSVIRSEAFSYEQALRIDDKTQND-----ELTMGMRAALLKLSVKLKQKRLLEAGA            |     |     |     |     |     |     |     |
| MM-LOBO        | 476 | EMFGRTIIRSCTKLSYDHAQSMIENPTEKIPPEELPPISPEHSVEEVHQAVLNHLSIAKQLRRQRFVDGA  |     |     |     |     |     |     |     |
| HS-LOBO        | 103 | EMFGRTIIRSCTKLSYHAQSMIESPTEKIPAKELPPISPEHSSEEVHQAVLNHLSIAKQLRRQRFVDGA   |     |     |     |     |     |     |     |
| CE-Q09568-LOBO | 415 | VWFGRSVIRSRVKLAYEHAQDFIENPEKDFTCDELPDISDGNTPFEIKEKTLMLHRIAQVLRQKREDSGA  |     |     |     |     |     |     |     |
| SP-Z99259      | 549 | RWFGKTVIKTCARLAYSEAQGVIEGKSWDDAVG--KPIGGTHTPKOVETSILTLCFISRKLKDKRFAKGA  |     |     |     |     |     |     |     |
| SC-P24276-SSD1 | 798 | TWVGESTISPSNLSLEQLDEKLSGTSPTS-----YLSVTQEIARSFYARRINDPE                 |     |     |     |     |     |     |     |
| SC-P39112      | 648 | PKIRKGIVSNFPKATYEDVDRILGTPNSEAS-----PVKKDLESLSMISKLLREQRIKNSN           |     |     |     |     |     |     |     |
|                |     | ----- ----- ----- ----- ----- ----- ----- ----- -----                   |     |     |     |     |     |     |     |
|                |     | 841   | 851 | 861 | 871 | 881 | 891 | 901 |     |

Fig. 2h

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|                |     |  |     |     |      |      |      |      |      |
|----------------|-----|--|-----|-----|------|------|------|------|------|
| HS-HMC-Dis3    | 628 | LTLSSPEVRFHMDSETHDPIDLQTKELRETNSMVEEFMLLANISVAKKIHIEEFSEHALLRKHPPPPSNY     | 911 | 921 | 931  | 941  | 951  | 961  | 971  |
| MM-hmc-Dis3    | 628 | LTLSSPEIRPHMDSETHDPIDLQTKELRETNSMVEEFMLLANISVAKKIHIEEFSEHALLRKHPPPPSNY     |     |     |      |      |      |      |      |
| CE-Q17632-Dis3 | 641 | LTLASSEVRFDMDWESRTPKKVMEKQHLDTHTSMVEEFMLLANISVAEKILEEYPCALLRRHPVPLKESY     |     |     |      |      |      |      |      |
| SP-P37202-Dis3 | 658 | LNLASPEVRIQTDNETSDPMDVEIKQLLETNSLVEEFMLLANISVAQKIYDAFPQTAVLRRHAAPPLTNF     |     |     |      |      |      |      |      |
| SC-Q08162-Dis3 | 693 | LNLASPEVKVHMDSETSDPNEVEIKLLATNSLVEEFMLLANISVARKIYDAFPQTAVLRRHAAPPLTNF      |     |     |      |      |      |      |      |
| MM-LOBO        | 546 | LRLDQLKLAFTLDHETGLPQGCCHIYEYRESNKLVEEFMLLANMAVAHKKIFRTTPEQALLRRHPPPPQTKML  |     |     |      |      |      |      |      |
| HS-LOBO        | 173 | LRLDQLKLAFTLDHETGLPQGCCHIYEYRESNKLVEEFMLLANMAVAHKKIHRAFPPEQALLRRHPPPPQTKML |     |     |      |      |      |      |      |
| CE-Q09568-LOBO | 485 | LRIELPRLKFALDEDK-KPQGVSIYEIKDSNKLVEEFMLLANMEVAKKIAENFPEHALLRNHPPPPKEKMI    |     |     |      |      |      |      |      |
| SP-Z99259      | 617 | VEINSTELKQLDSEYG-MPNKCEVYEQTDANHLIEEFMLLANRSVAEHI SKNFSNNSLLRRHASPKEKQI    |     |     |      |      |      |      |      |
| SC-P24276-SSD1 | 849 | ATLLPTLSLLESLEDEKVKVDINILDRTLGFVINEIKRKVNSTVAEKIYTKLGLDLALLRRROMQPIATKM    |     |     |      |      |      |      |      |
| SC-P39112      | 704 | AVIFGEGFNKGLVMIN---ADSEGELTEVTFSDQEEITLSTILVSEMMILANTLTGRYPAEN---KI        |     |     |      |      |      |      |      |
|                |     | ----- ----- ----- ----- ----- ----- ----- ----- ----- -----                | 911 | 921 | 931  | 941  | 951  | 961  | 971  |
| HS-HMC-Dis3    | 698 | EILVKAARS-RNLEIKTDTAKSLAESLDQAESPTFPYLN--TLRLILATRCMMQAVYFCSGMD-----       |     |     |      |      |      |      |      |
| MM-hmc-Dis3    | 698 | DILVKAARS-KNLQIKTDTAKSLADSLDRAESPDFFYLN--TLRLILATRCMMQAVYFCSGMD-----       |     |     |      |      |      |      |      |
| CE-Q17632-Dis3 | 711 | KPLVEAARH-RGFEIIVESGKGLADSLNRCVDKKNPMLN--RLRLMLTTRCMTQAVYFSAKDGFSDIKL      |     |     |      |      |      |      |      |
| SP-P37202-Dis3 | 728 | DSLQDIILRVCKGMHLKCDTSKSLAKSLDECVDPREPYFN--TLRLILATRCMLSAEYFCSGTFAP-----    |     |     |      |      |      |      |      |
| SC-Q08162-Dis3 | 763 | EILNEMLANTRKNMSISLESSKALADSLDRCDPEDPYFN--TLVRIMSTRCMMAAQYFYSGAYS-----      |     |     |      |      |      |      |      |
| MM-LOBO        | 616 | SDLVEFCDO-MGLPMDVSSAGALNKSLTKTFGDDKYSLARKEVLTNNYSRPMQMALYFCSGMLQD-----     |     |     |      |      |      |      |      |
| HS-LOBO        | 243 | SDLVEFCDO-MGLPMDVSSAGALNKSLTKTFGDDKYSLARKEVLTNNYSRPMQMALYFCSGMLQD-----     |     |     |      |      |      |      |      |
| CE-Q09568-LOBO | 554 | KDVAEQCAR-IGFPIDGRTSGLLSTSLRKYQKSLDMCIRQVSSLTIKPMQAKYFCT--FEM-----         |     |     |      |      |      |      |      |
| SP-Z99259      | 686 | NEFCHFLKS-MNFDFFDASSAAFNASMVRLRSTNRELV--ELFENMAVRSINRAEYFCTGDFGEK-----     |     |     |      |      |      |      |      |
| SC-P24276-SSD1 | 919 | ASFRKKIQN-FGYNFDTNTADELIKGVLIKDDDDVRVGI-----EILFKTWPRAKRYFIAGKVD-----      |     |     |      |      |      |      |      |
| SC-P39112      | 764 | GGVFRCYKQ---LPLGEVAQQQYDSMITSTKKGITPKLK-----DIVKLSLLNSSFTGRPFR-----        |     |     |      |      |      |      |      |
|                |     | ----- ----- ----- ----- ----- ----- ----- ----- ----- -----                | 981 | 991 | 1001 | 1011 | 1021 | 1031 | 1041 |

Fig. 2i

[illegible]

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|                |     |          |  |
|----------------|-----|----------|--|
| HS-HMC-Dis3    | 835 | QLFFKSKG | IVSEEAYILFVRKNAIVVLIPKYGLEGTVFPEEKDKPNPQLI |
| MM-hmc-Dis3    | 835 | QLFFKSKG | IVSEEAYILFVRKNAIVVLIPKYGLEGTVFPEEKDKPKPRLA |
| CE-Q17632-Dis3 | 918 | VRYFKGK  | VETCEGFVNGVRNNGIQVFPKYGLESILVLQTSASG--TI   |
| SP-P37202-Dis3 | 868 | GQALKGG  | VAEEDAYVIXVFXNGFVVFIAFGLGEGIVYTKSLSSVLEPN  |
| SC-Q08162-Dis3 | 903 | GQVMRNN  | ESTETGYVIXVFXNGFVVFIAFGLGEGIVYTKSLSSVLEPN  |
| MM-LOBO        | 756 | AVLVKESG | PLESEAMVNGVLNQAFDVLVLRFGVQKRIYCNALALRSYSFQ |
| HS-LOBO        | 383 | AVLVKESG | PLESEAMVNGILKQAFDVLVLRFGVQKRIYCNALALRSYHFQ |
| CE-Q09568-LOBO | 693 | GVFIHQTG | PMKCQAVVLGVMDLSFDVLIVEYGVVVRVVDKMR--DEN    |

Fig. 2k

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|                |      |  |      |      |      |      |      |      |      |
|----------------|------|--|------|------|------|------|------|------|------|
| SP-Z99259      | 823  | SVYIAEYCKKHDKXMPVQAFATRISGNSIDVYISEYGISNRVDLSSDDR-IKSP1-----           | 1191 | 1201 | 1211 | 1221 | 1231 | 1241 | 1251 |
| SC-P24276-SSD1 | 1052 | CKTINDMGNTTG--QLLTMATVLQVYESSFDVFIPEFGIEKRVHGDQLPLIKAFFDGTNRVLELHMOPGV |      |      |      |      |      |      |      |
| SC-P39112      | 890  | LNVLKKL-----TKLEPERTFDVMT-----SVPQNGFTGCVFPDLSFARGTLK-----             |      |      |      |      |      |      |      |
|                |      |  | 1191 | 1201 | 1211 | 1221 | 1231 | 1241 | 1251 |
| HS-HMC-Dis3    | 885  | -----YDDEIPSLKIED-TVFHVFDKVKVKIMLDSSNLQHOKIRMSLVEPQIPGIS               |      |      |      |      |      |      |      |
| MM-hmc-Dis3    | 885  | -----YDDEIPSLRIEG-TVFHVFDKVKVKITLDSSNLQHOKIRMALVEPQIPGIN               |      |      |      |      |      |      |      |
| CE-Q17632-Dis3 | 965  | -----IDVEEMSVKVGWDVWIKELFPTVRISVNEKNQQRPRVELQLIKPAIPGLS                |      |      |      |      |      |      |      |
| SP-P37202-Dis3 | 916  | -----VEVVEDEYKINI--EIRDQPKPQT---VQIQMFQQVVRVTTVRDEHSGKQ                |      |      |      |      |      |      |      |
| SC-Q08162-Dis3 | 950  | -----AAFDEVEYKLTFF--VPTNSDKPR-----DVVVFQVVEVQVRSVMDPITSKR              |      |      |      |      |      |      |      |
| MM-LOBO        | 806  | -----KVGKKPELTAVWEPD-DLE---EPTQQQVITIPSLVDVWLQAEATALKYS                |      |      |      |      |      |      |      |
| HS-LOBO        | 433  | -----KVGKKPELTAVWEPE-DME---QEPAAQQVITIPSLVEVWLQAEATALKYS               |      |      |      |      |      |      |      |
| CE-Q09568-LOBO | 740  | -----KSTEK--LTIVPADPNAESGNREEFSSSIQMCNVVYVIL-VPIKXSIEVSA               |      |      |      |      |      |      |      |
| SP-Z99259      | 878  | -----VAPDDSSVKITL---FDDSS---Q-K---TIALTDRFQVYLYSDYSRTFFSI              |      |      |      |      |      |      |      |
| SC-P24276-SSD1 | 1120 | DSATFIPADEKNPKSYRNSIKNKRSTAAEIANIELDKRAESEPLISDPLSKELSDLHILTVPNRLPSAS  |      |      |      |      |      |      |      |
| SC-P39112      | 934  | -----LHPSSMHYPMIG-----DIVKNCKISKIDCLEGMLELEKL-----                     |      |      |      |      |      |      |      |
|                |      |  | 1261 | 1271 | 1281 | 1291 | 1301 | 1311 | 1321 |

Fig. 21

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|                |      |             |   |      |      |      |      |
|----------------|------|-------------|---|------|------|------|------|
| HS-HMC-Dis3    | 935  | IPTDTS      | -----NMDLNGPKKKKKMLGK-----                        |      |      |      |      |
| MM-hmc-Dis3    | 935  | IPPNVA      | -----DKALTAPGKKRRKLEK-----                        |      |      |      |      |
| CE-Q17632-Dis3 | 1016 | V           | -----DFDLSSEG-----LGL-----                        |      |      |      |      |
| SP-P37202-Dis3 | 962  | K           | -----V-----QITLVY-----                            |      |      |      |      |
| SC-Q08162-Dis3 | 994  | K           | -----A-----ELLLK-----                             |      |      |      |      |
| MM-LOBO        | 853  | ILKRPQ      | -----LEKASDEEPED-----                             |      |      |      |      |
| HS-LOBO        | 480  | ILKRPQTQGH  | LGPEKEEESDGEPEDSSTS                               |      |      |      |      |
| CE-Q09568-LOBO | 788  | TIVRPS      | -----LEQRNLIKSTLKDMMKETGSTILQ-----                |      |      |      |      |
| SP-Z99259      | 919  | R           | -----CSLVSLN-----                                 |      |      |      |      |
| SC-P24276-SSDI | 1190 | DNKQNALEKFI | STTETRIENDNYIQEIHELQKIPILLRAEVGMALPCLTVRAINPFMKRV |      |      |      |      |
|                |      |             | ----- ----- ----- ----- ----- ----- ----- -----   |      |      |      |      |
|                |      |             | 1331  | 1341 | 1351 | 1361 | 1371 |
|                |      |             |   |      |      |      | 1381 |

Fig. 2m

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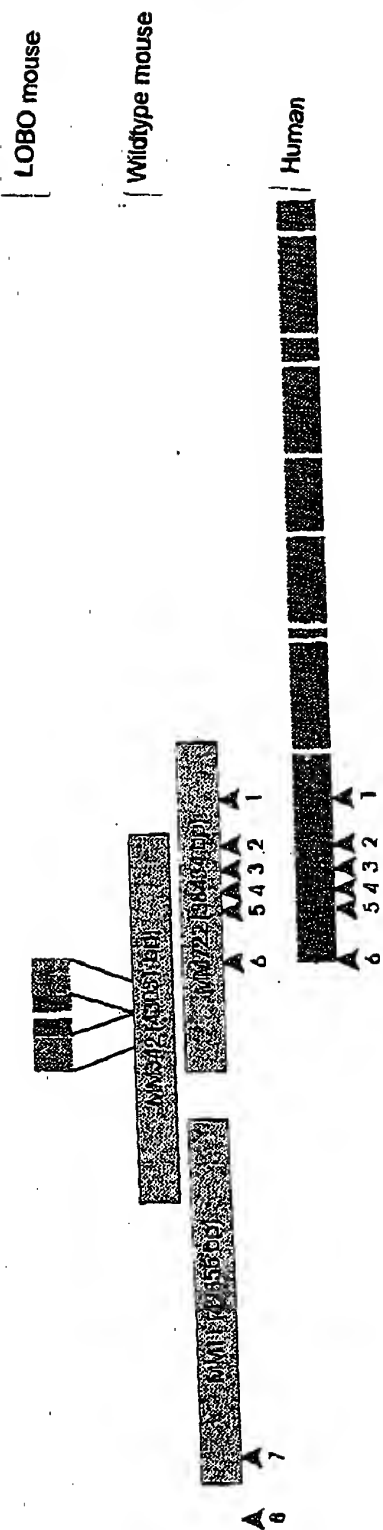


Fig. 3

WT LOBO

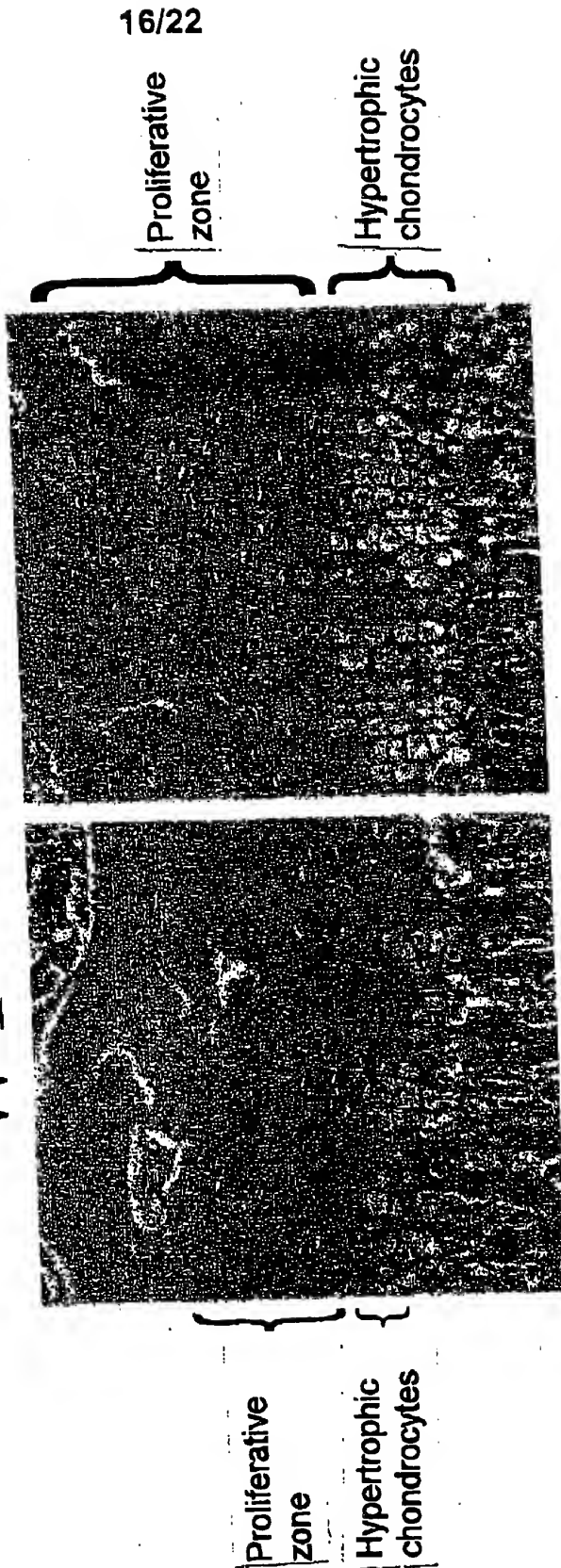


Fig. 4



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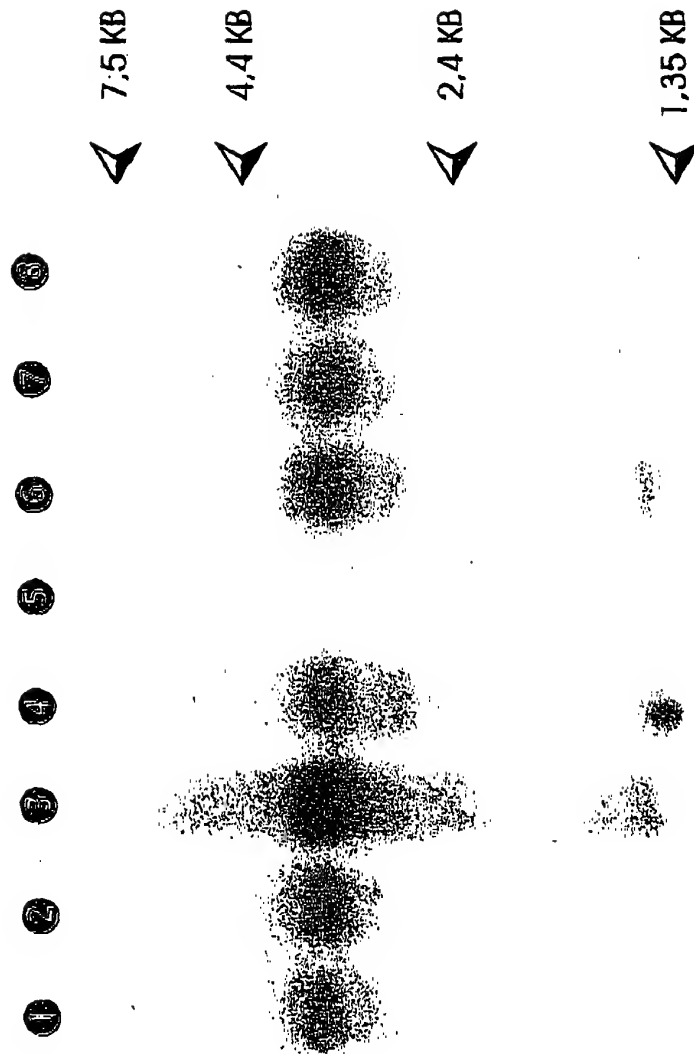


Fig. 5

# Dis3 Proteins

## VACB / RNase II

## LOBO

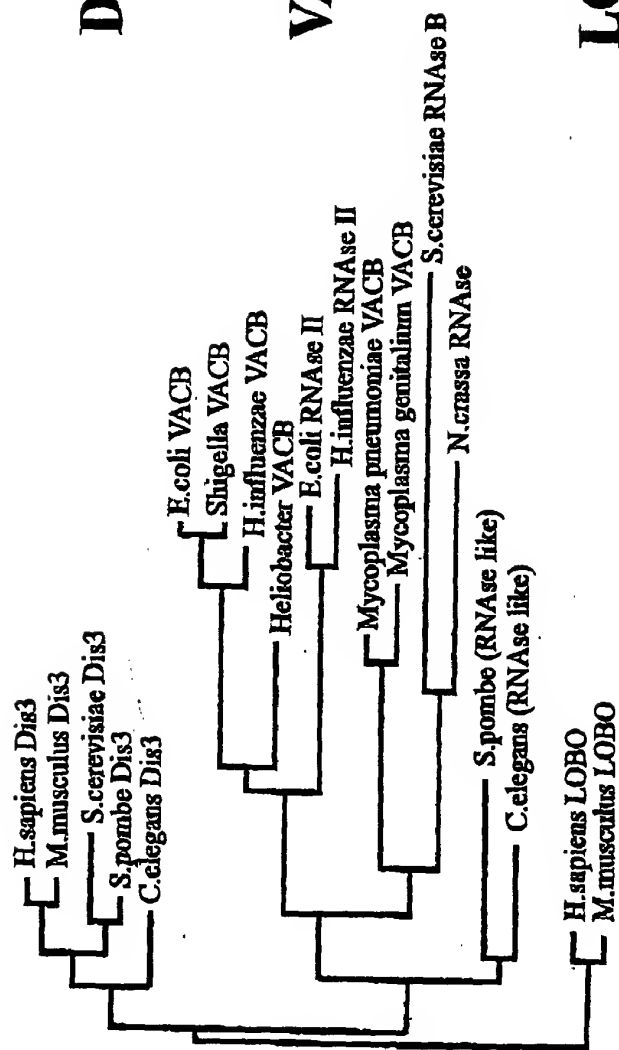


Fig. 6

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LOBO



WT



Fig. 7

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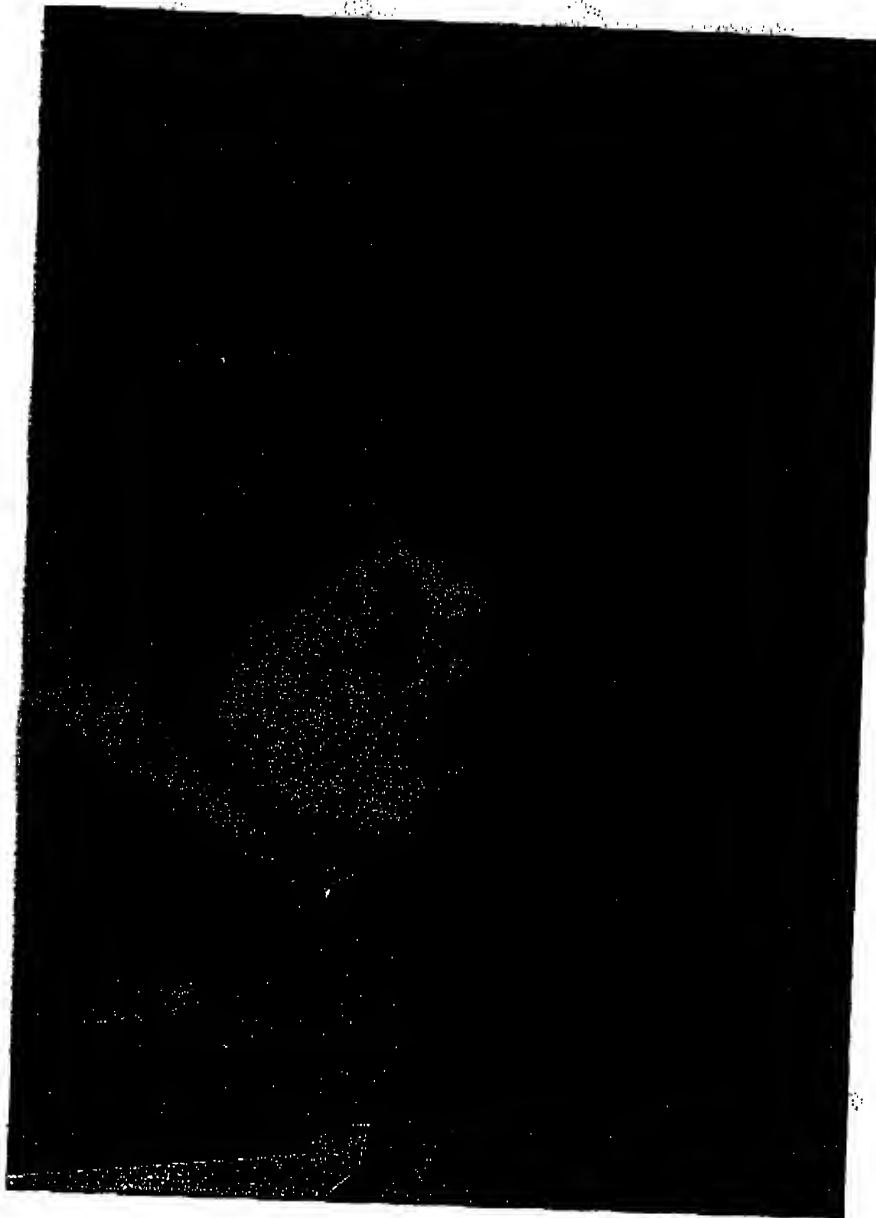


Fig. 8

Fig. 8  
The figure shows a dark, rectangular area, possibly a photograph or a heavily underexposed image, occupying the central portion of the page.

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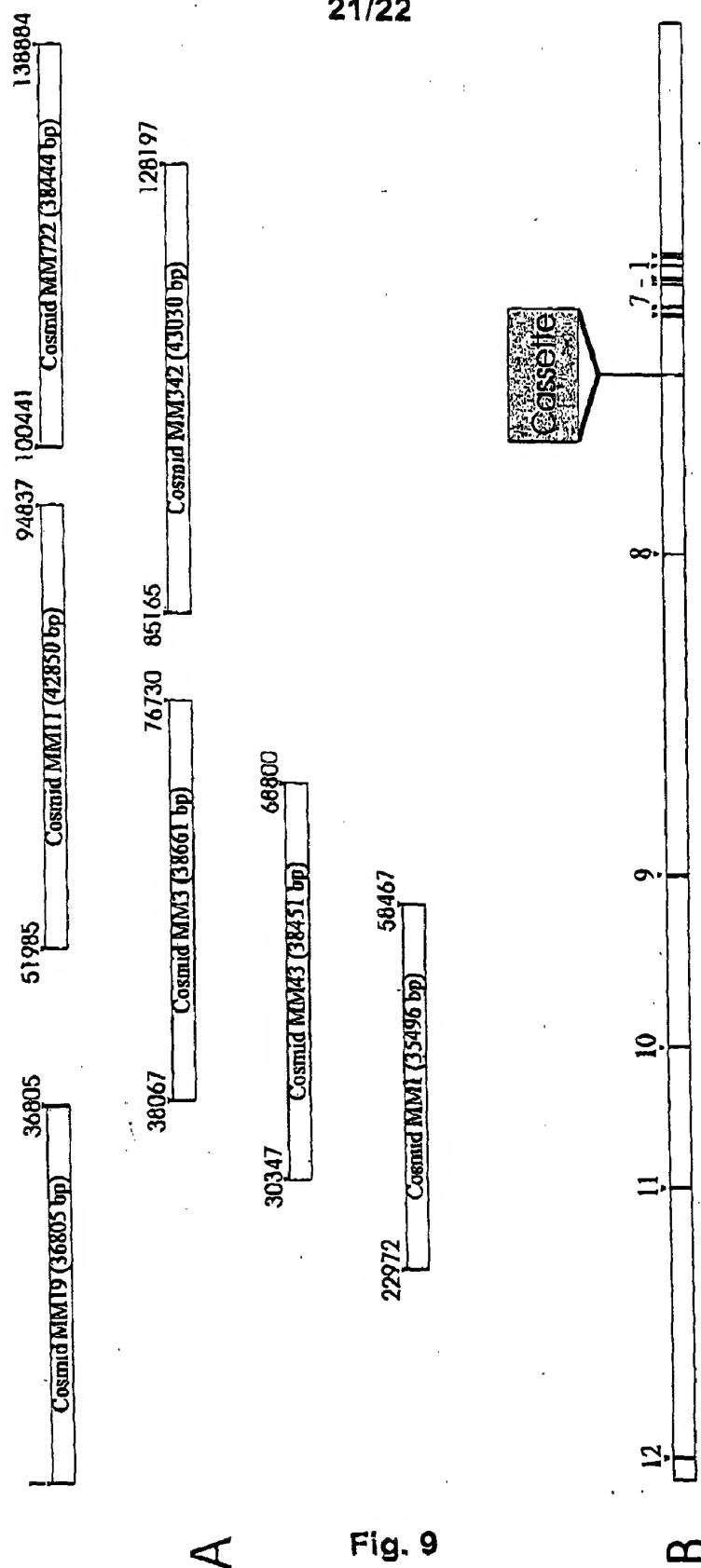


Fig. 9

STS Marker  
WI-8964

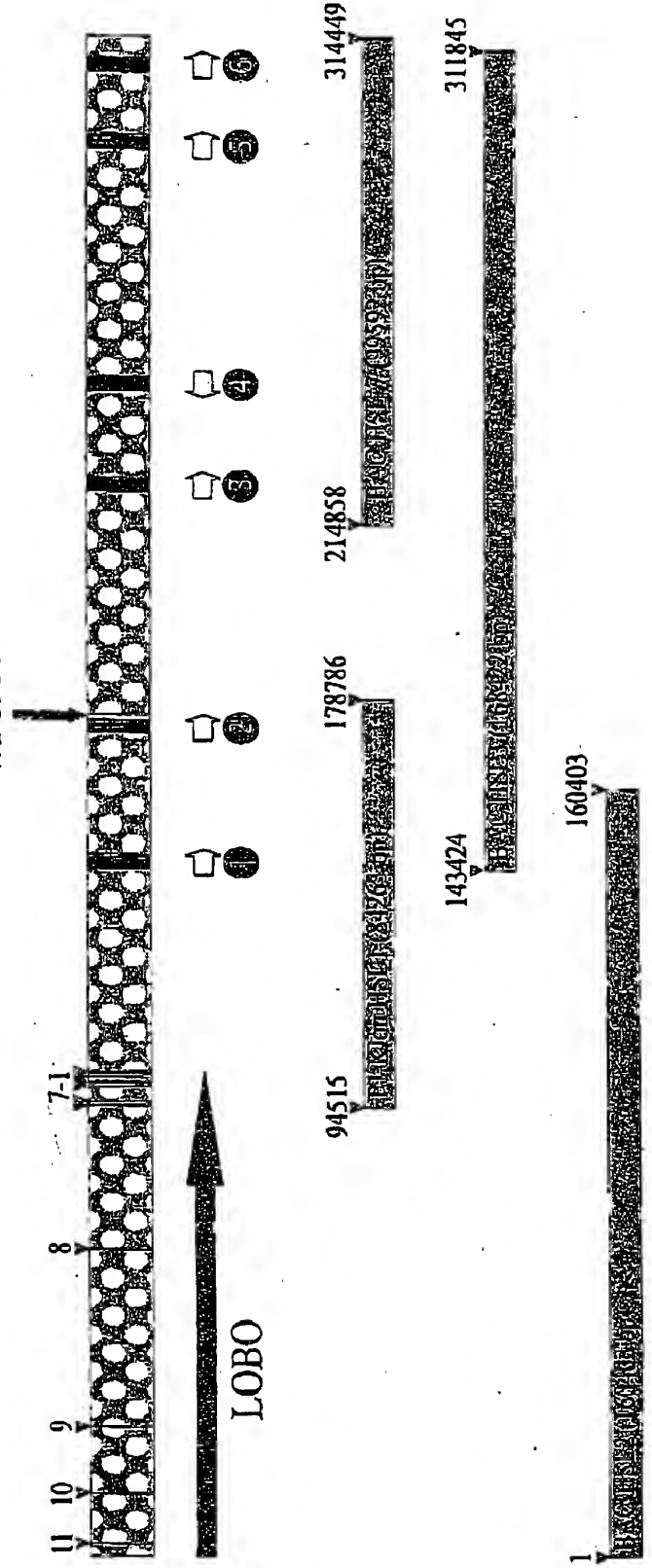


Fig. 10